

1600

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/486,676

DATE: 01/08/2002

TIME: 15:22:34

Input Set : A:\GJE-39.ST25.txt

Output Set: N:\CRF3\01082002\I486676.raw

ENTERED

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3 <110> APPLICANT: Crisanti, Andrea
 5 <120> TITLE OF INVENTION: Conjugates that Contain the Homeodomain of Antennapedia
 7 <130> FILE REFERENCE: GJE-39
 9 <140> CURRENT APPLICATION NUMBER: US 09/486,676
10 <141> CURRENT FILING DATE: 2000-03-01
12 <150> PRIOR APPLICATION NUMBER: GB 9718609.2
13 <151> PRIOR FILING DATE: 1997-02-02
15 <160> NUMBER OF SEQ ID NOS: 12
17 <170> SOFTWARE: PatentIn version 3.1
19 <210> SEQ ID NO: 1
20 <211> LENGTH: 60
21 <212> TYPE: PRT
22 <213> ORGANISM: Drosophila sp.
24 <400> SEQUENCE: 1
26 Arg Lys Arg Gly Arg Gln Thr Tyr Thr Arg Tyr Gln Thr Leu Glu Leu
                                       10
30 Glu Lys Glu Phe His Phe Asn Arg Tyr Leu Thr Arg Arg Arg Ile
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34 Glu Ile Ala His Ala Leu Cys Leu Thr Glu Arg Gln Ile Lys Ile Trp
                               40
38 Phe Gln Asn Arg Arg Met Lys Trp Lys Lys Glu Asn
42 <210> SEQ ID NO: 2
43 <211> LENGTH: 8
44 <212> TYPE: PRT
45 <213> ORGANISM: Artificial Sequence
47 <220> FEATURE:
48 <223> OTHER INFORMATION: Six histidine tail fused to conjugate for purification
50 <400> SEQUENCE: 2
52 His His His His His Gly Ser
53 1
56 <210> SEQ ID NO: 3
57 <211> LENGTH: 16
58 <212> TYPE: PRT
59 <213> ORGANISM: Drosophila sp.
61 <400> SEQUENCE: 3
63 Arg Gln Ile Lys Ile Trp Phe Gln Asn Arg Arg Met Lys Trp Lys Lys
64 1
                                       10
67 <210> SEQ ID NO: 4
68 <211> LENGTH: 16
69 <212> TYPE: PRT
70 <213> ORGANISM: Artificial Sequence
72 <220> FEATURE:
73 <223> OTHER INFORMATION: Variant sequence of helix 3. Must have between
         6 and 10 hydrophobic amino acids.
76 <220> FEATURE:
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77 <221> NAME/KEY: MISC_FEATURE

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Input Set : A:\GJE-39.ST25.txt Output Set: N:\CRF3\01082002\I486676.raw 78 <222> LOCATION: (1)..(5) 79 <223> OTHER INFORMATION: Can be any alpha-amino acid. 82 <220> FEATURE: 83 <221> NAME/KEY: MISC_FEATURE 84 <222> LOCATION: (7)..(16) 85 <223> OTHER INFORMATION: Can be any alpha-amino acid. 88 <400> SEQUENCE: 4 90 Xaa Xaa Xaa Xaa Xaa Trp Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa 91 1 94 <210> SEQ ID NO: 5 95 <211> LENGTH: 16 96 <212> TYPE: PRT 97 <213> ORGANISM: Artificial Sequence 99 <220> FEATURE: 100 <223> OTHER INFORMATION: Variant sequence of helix 3. Must have between 6 and 10 hydrophobic amino acids. 103 <220> FEATURE: 104 <221> NAME/KEY: MISC_FEATURE 105 <222> LOCATION: (1)..(10) 106 <223> OTHER INFORMATION: Can be any alpha-amino acid. 109 <220> FEATURE: 110 <221> NAME/KEY: MISC_FEATURE 111 <222> LOCATION: (12)..(16) 112 <223> OTHER INFORMATION: Can be any alpha-amino acid. / 115 <400> SEQUENCE: 5 117 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Trp Xaa Xaa Xaa Xaa 118 1 121 <210> SEQ ID NO: 6 122 <211> LENGTH: 35 123 <212> TYPE: PRT 124 <213> ORGANISM: Drosophila sp. 126 <400> SEQUENCE: 6 128 Leu Thr Arg Arg Arg Ile Glu Ile Ala His Ala Leu Cys Leu Thr 132 Glu Arg Gln Ile Lys Ile Trp Phe Gln Asn Arg Arg Met Lys Trp Lys 20 136 Lys Glu Asn 137 35 140 <210> SEQ ID NO: 7 141 <211> LENGTH: 35 142 <212> TYPE: PRT 143 <213> ORGANISM: Drosophila sp. 145 <400> SEQUENCE: 7 147 Leu Thr Arg Arg Arg Ile Glu Ile Ala Tyr Ala Leu Cys Leu Thr 10 151 Glu Arg Gln Ile Lys Ile Trp Phe Ala Asn Arg Arg Met Lys Trp Lys 25 155 Lys Glu Asn

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156

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159 <210> SEQ ID NO: 8
160 <211> LENGTH: 35
161 <212> TYPE: PRT
162 <213> ORGANISM: Drosophila sp.
164 <400> SEQUENCE: 8
166 Leu Thr Arg Arg Arg Ile Glu Ile Ala His Ala Leu Cys Pro Pro
170 Glu Arg Gln Ile Lys Ile Trp Phe Gln Asn Arg Arg Met Lys Trp Lys
                                     25
171
174 Lys Glu Asn
            35
175
178 <210> SEQ ID NO: 9
179 <211> LENGTH: 4
180 <212> TYPE: PRT
181 <213> ORGANISM: Artificial Sequence
183 <220> FEATURE:
184 <223> OTHER INFORMATION: Factor Xa cleavage sequence.
186 <400> SEQUENCE: 9
188 Ile Glu Gly Arg
189 1
192 <210> SEQ ID NO: 10
193 <211> LENGTH: 5
194 <212> TYPE: PRT
195 <213> ORGANISM: Artificial Sequence
197 <220> FEATURE:
198 <223> OTHER INFORMATION: Enterokinase cleavage sequence.
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202 Asp Asp Asp Asp Lys
203 1
206 <210> SEQ ID NO: 11
207 <211> LENGTH: 5
208 <212> TYPE: PRT
209 <213> ORGANISM: Artificial Sequence
211 <220> FEATURE:
212 <223> OTHER INFORMATION: Thrombin cleavage sequence.
214 <400> SEQUENCE: 11
216 Leu Val Pro Arg Gly
217 1
220 <210> SEQ ID NO: 12
221 <211> LENGTH: 17
222 <212> TYPE: DNA
223 <213> ORGANISM: Artificial Sequence
225 <220> FEATURE:
226 <223> OTHER INFORMATION: GAL4 DNA binding domain.
228 <400> SEQUENCE: 12
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229 cggaggacag tcctccg

17

VERIFICATION SUMMARY

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L:90 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 L:117 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5